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ElogD_{oct}: A Tool for Lipophilicity Determination in Drug Discovery. Basic and Neutral Compounds

Field of the Invention

This invention is related to an improved method for determination of ElogD_{oct} values for drug candidates.

Background of the Invention

The importance of lipophilicity can be understood, for example, by considering the correlation between high lipophilicity and poor solubility, which has generally been explored with neutral solutes. Furthermore, lipophilicity has been shown to be of paramount importance in several other Absorption, Distribution, Metabolism, Excretion (ADME) aspects, that is, absorption, distribution, metabolism and excretion. It is generally held that very lipophilic compounds are "preferred" targets for metabolism, often leading to high clearance values and, quite often, lipophilicity positively correlates with a high plasma protein binding. A large volume of distribution, probably due to a high fraction of the compound bound to tissues, is often observed for lipophilic compounds. At physiological pH many basic or acidic drugs are ionized, and the partition coefficient is indeed a distribution coefficient, D, which is generally taken to be the distribution between an aqueous buffer at pH 7.4 and n-octanol, and it is indicated by the notation $D_{oct}^{7.4}$.

Scherrer³ defines the distribution coefficient (in the form of its logarithm) for monoprotic bases as:

$$\log D_{oct} = \log P_{oct} + \log \left[1 / (1 + 10^{pKa - pH}) \right]$$

For monoprotic acids the equation has the same form, except that the exponent is written as pH-pKa. For polyprotic compounds the equations become more complicated, and these aspects have been described in detail by Avdeef.⁶

Given the widespread use and application of distribution coefficients, a method that can accurately and rapidly yield $\log D_{oct}^{7.4}$ values, would be a welcome addition to the experimental tools available for physicochemical properties screening in the discovery setting. The general notation $\log D_{oct}$, is defined herein, to mean the values determined at pH 7.4.

Although computational packages for the estimation of logD_{oct} are available,⁷ or this value could also be calculated from estimated logP_{oct} and pKa values, we find that, for drug molecules, computed values are often inaccurate. Depending on the software used, they may differ by as much as two to three logD_{oct} units, among different software packages and/or from experimental values, since the accuracy of pKa as well as logP_{oct} has to be factored in. These methods are, of course, valuable when virtual libraries (or individual virtual molecules) are being designed and, with proper training, more accurate values might be obtained. However, as early as possible, and especially if a compound-sparing method is available, the

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computed values should be replaced by measured values, with particular regards to cases where intramolecular H-bonding is possible, and/or in the presence of conformational flexibility, and/or with molecules which can tautomerize. These occurrences typically offer an even greater challenge to fragment-based software packages. Structure Activity Relationships (SAR) and Structure Property Relationships (SPR) analyses and alerts such as the Lipinski "rule of 5",8 would greatly benefit by the introduction of accurate experimental values.

The classical shake-flask method, or variations of this method which have been described, are neither rugged nor rapid enough for medium to high-throughput applications, they are generally more sensitive to impurities, less amenable to automation than are RP-HPLC methods, and they do not usually offer a wide dynamic range.

RP-HPLC retention data, expressed either as $\log k$ or $\log k_w$, have been shown to correlate well with absolute and relative lipophilicity values to but, they have also been criticized as not being a true "replacement" for shake-flask values. Part of the criticism may stem from the limited scope of some reports, focusing either on fairly simple monofunctional solutes, or classes of analogs with a limited $\log P_{\rm oct}$ range, and in many cases there was only mention of neutral solutes, under the pH conditions of the method. When these correlations were extended beyond classes of analogs, less encouraging results were obtained. Furthermore, in several cases, the slope of these correlations was quite different from unity, casting doubts about the different balance of forces responsible for the two values. Indeed, LFER (Linear Free Energy Relationships) analyses have shown that $\log k$ on typical RP-HPLC systems do not encode the same blend of factors, as does $\log P_{\rm oct}$. In particular, $\log k$ values respond to solute hydrogen-bond acidity, but $\log P_{\rm oct}$ values do not.

k represents the capacity factor of the solute at a given organic solvent concentration, and k_w is the capacity factor extrapolated to a 0% concentration of the organic solvent. The observations reported above pointed us toward using the extrapolated value rather than, as reported by Yamagami and co-workers, 15 a log k value, which is likely to be limited in its applicability to a wide variety of drug-like compounds (amphiprotic) and, therefore, of limited use for the goals we set. As we reported previously 16, a judicious choice of columns and mobile phase, as well as flow rates, would greatly enhance the performance of RP-HPLC methods.

Another factor of great importance, if the data were to be used for software training purposes, or for the creation of a large database, is the reproducibility from column to column, which depends on the quality of the packing chemistry and manufacturing. We have shown, in the case of the ElogP data, ¹⁶ that the latter aspect does not appear to be a problem.

However, it is advisable to monitor the column performance for possible deterioration, especially if high-throughput screening is the goal.

The speed of the determination and the ability to handle diverse structures and lipophilicity values are, of course, of paramount importance in an industrial research setting. These aspects translate into the capability of screening, with modest resources, a large number of compounds, with a good degree of accuracy across a wide range of lipophilicity values and hydrogen-bonding properties.

Summary of the Invention

This invention provides a method that is accurate, rapid, and possesses a good dynamic range, together with being applicable to a variety of drug-like molecules, and which is robust with respect to ion-pairing and concentration-related variability.

Although most of the determination of $LogD_{oct}$ disclosed herein were conducted at pH 7.4, other buffers with pH values ranging from pH 4 to 8 may be employed advantageously at pH values to mimic the physiological values in different parts of the body where different drugs are functional or absorbed in the blood stream.

This invention provides a method of experimentally determining $ElogD_{oct}$ for chemical compounds (" $ElogD_{oct}$ " is the notation herein for "experimentally determined" $logD_{oct}$, with " $logD_{oct}$ ", unless otherwise indicated, referring to the classical shake-flask or literature value) which comprises:

- (a) Introducing said chemical compounds seriatim to the column of a reverse phase high performance liquid chromatographic system said column being an embedded amide functional group column; or a C-18 bonded column with low silanol activity; and
- (b) Eluting said compounds with a mobile phase containing Morpholine Propane Sulfonic (MOPS) buffer and a methanol/octanol mixture in which the proportions of said methanol/octanol mixture to said buffer are from 75 to 15% v/v; and with flow rates between 0.5 and 3 ml/min and
 - (c) Measuring the retention time required to elute each sample from said column; and
- (d) Calculating ElogD_{oct} from the retention time of each sample using equation1 (the derivation of which will be discussed).

This invention further provides a method wherein said compounds for which $ElogD_{oct}$ is to be determined are divided into groups according to calculated lipophilicity based on chemical structure and; $ElogD_{oct}$ is determined for all samples in a first group and; said column is equilibrated to the conditions for a second group.

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This invention further provides a method wherein said $logD_{oct}$ values are calculated by a programmed computer and samples within a defined $logD_{oct}$ range are introduced seriatim into said column by robotic means under control of a programmed computer and calculating and recording each $ElogD_{oct}$ from the retention time of each sample by equation 1.

This invention further provides a method wherein each of steps a) through d) is performed by robotic means under the control of a programmed computer.

This invention further provides a method wherein said column is an embedded amide functional group column.

This invention further provides a method wherein said column is a C-18 bonded column with low silanol activity.

This invention further provides a method wherein logD_{oct} may be determined at a buffer pH of 4 to 8 and preferably at 6.5.

These and other features and advantages of the present invention will become more evident from the following discussion and drawings wherein:

Short Description of the Drawings

Figure 1. is a correlation between $\log D_{\rm oct}$ and $\log k_w^{'}$ for 90 solutes; and Figure 2 is a plot of residuals vs. $\log D_{\rm oct}$

Detailed Description of the Invention, Preferred Embodiment and Drawings

For a subset of the compounds, including basic and neutral compounds, an addition of *n*-decylamine to the mobile phase yields improved correlations, although it was not necessary for the neutral solutes. ¹⁶ In order to maximize the speed of analysis, while still retaining a good accuracy, the same flow rate previously reported ¹⁶ was chosen for each range of lipophilicity. We also studied the effect of a further increase of flow rate on accuracy, and performed statistical analysis in order to determine experimental lipophilicity "thresholds" for each range of conditions, which would yield accurate results.

Extensive work was conducted using the 90 solutes reported in Table 1, and each $\log k_{\rm w}$ value is the average of at least three determinations, on different columns, with an average standard deviation of 0.07. The table also reports the standard deviation of $\log k_{\rm w}$ for each compound (N =3 to 5) and no large deviations were observed, regardless of structure and range of lipophilicity, with methotrimeprazine yielding the largest s.d. of 0.27. It is worth noting that routine potentiometric lipophilicity determinations have a typical s.d. of 0.4 $\log D_{\rm oct}$ units, for replicate determinations, and often they have to be extrapolated to 0% organic solvent from mixed solvents, due to the poor aqueous solubility of many compounds.

The fit of the $\log k_w$ to the averaged $\log D_{\rm oct}$ values is reported below (Eq.1), and shown graphically in Figure 1.

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 $log D_{oct} = 1.1267 (\pm 0.0233) log k'_{w} + 0.2075 (\pm 0.0430)$ (Eq. 1)

N = 90, $R^2 = 0.964$, R = 0.982, s = 0.309, F = 2339, $q^2 = 0.962$ (these values are standards related to statistical confirmation of the derived values, with N being the number of compounds, R being the correlation coefficient (R^2 is the square thereof), s being the standard deviation, F being the Fisher statistic and q^2 being calculation deviation with sequential calculation repetitions with omission of one compound).

The slope obtained is very close to unity, with a small intercept, and these parameters offer a good comparison of the balance of forces which play a role in the shakeflask vs. RP-HPLC distribution coefficient. The question of the diagnostic importance of the slope has been stressed by Minick et al. 17 Pointing to the work of Melander et al., 20 these authors state that "...equations correlating $\log k_{\rm w}$ and $\log P_{\rm oct}$ data represent linear free energy relationships in which the slope is an estimate of how closely the free energies of the processes compare." A slope close to unity implies that the two processes are homoenergetic, i.e. the free energy changes are the same. Furthermore, if the goal is the determination of the "classical" logDoct, then a slope significantly different from one would enhance any error in the determination of or logDoct. And a slope significantly different from unity is an indication of a fairly large over- or underestimation of lipophilicity by the method. Obviously, if a different scale of lipophilicity is the goal, $\log k_w$ values could be used as such, or different indices could be developed. Valkó et al., 21 described a chromatographic hydrophobicity index (CHI), obtained via a gradient run. In this case a correlation with a "classical" shake flask logPoct was not necessarily sought, and a self-consistent CHI scale was established. However, logDoct data are so widely used in many correlations by the medicinal chemistry community, that a "classical" logDoct value is likely to be desired. To the best of our knowledge no other method, capable of encompassing all the accuracy and ruggedness requirements we set as goals for this work, including a very practical set of conditions and speed, has been reported in the literature to date. By analogy with our previous work¹⁶ we termed the values obtained via Eq. 1 as ElogD_{oct}, and we will refer to them as such hereinafter.

A plot of residuals vs. $logD_{oct}$ values, as in Figure 2, shows that the error distribution is very consistent across the entire range, and no curvature (larger error) is observed at extreme values. This is important because it shows that similarly accurate determinations can be obtained across a dynamic range of 7 $logD_{oct}$ units, and the standard error is fairly small, considering also the variability of some of the $logD_{oct}$ data reported in the literature.

The question might be asked, about whether or not decylamine acts as a modifier other than a masking agent for potentially ionized silanols, since its absence is detrimental to the performance of the method. A comparison between $\log k_w$ values obtained under the

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conditions reported in our previous work 36 neutral solutes, ¹⁶ termed here $\log k_w$ (P), and the values obtained under the present condition for the same solutes, i.e. $\log k_w$ (D), shows that the balance of forces is the same, as demonstrated by a very small intercept (non-significant) and a slope very close to unity, in Equation 2. Since we have demonstrated that $\log k_w$ under our previous conditions encodes the same balance of forces as in a biphasic octanol-water system, ¹⁶ we conclude that the values obtained by this method are "true" $\log D_{oct}$ values.

$$\log k_w(P) = 1.0429 (\pm 0.0241) \log k_w(D) - 0.0219 (\pm 0.0522)$$
 (Eq. 2)
 $N = 36$, $R^2 = 0.982$, $R = 0.991$, $S = 0.198$, $S = 0.198$, $S = 0.980$

We have also tested a set of 10 proprietary compounds, basic and neutral, with molecular weights between 209 and 532 dalton. These compounds are structurally dissimilar from the compounds in the training set, and possess a wide range of functional groups. The results of the ElogD determinations are compared, in Table 2, with shake-vial and/or potentiometric determinations, and they show a good performance of the method for compounds 1-10.

As in our previous method, ¹⁶ we used control charts, as measure for a day-to-day system suitability check, which were constructed for ten compounds chosen across the entire range, and which are indicated in Table 1 (see Statistical Analysis section). An unexpected variation in these plots would immediately "flag" questionable results. Furthermore, we find that triflupromazine (CAS no. 146-54-3) offers a very sensitive "probe" for the column performance monitoring, as well as its analog chlorpromazine (CAS no. 50-53-3). A decline in the $\log k_w$ value of either of these two compounds is a good indication of column deterioration. A $\log k_w$ value for triflupromazine below 2.7 (2.6 for chlorpromazine) would indicate that the column should be replaced.

Currently, the ElogD determinations are run with the aqueous portion of the mobile phase prepared by a commercial laboratory, according to our specifications, and in fairly large batches (> 20 gallons), and we have noticed no significant difference in performance, after an initial adjustment of pH as needed. This practical enhancement helps with the speed of analysis, and it may also be taken as an indication of the ruggedness of the method.

As a further improvement, in terms of speed of analysis, we have also attempted to increase the flow rate and we have further automated the calculation procedure, through inhouse software, to obtain the final $ElogD_{oct}$ value, with very limited manual intervention, directly from the chromatographic data file. These modifications allow for an enhanced throughput, starting with an already rapid procedure. $ElogD_{oct}$ data for any compound are obtained, on average, in \leq 20 minutes, on a single instrument. Equation 3 shows a high correlation between the $log k_w^i$ values obtained with the "standard" flow rate (sf, 1.0 mL/min)

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and the faster flow rates (ff, 1.5 mL/min), used for a "mixed" set of data comprising 56 proprietary and commercial compounds, using the same compounds under each condition, and encompassing roughly 3 $\log k_w$ units, largely across the medium range defined in the Experimental section. We have run over 2,000 compounds with the "standard" flow rates.

$$\log k_w^{'}(\text{sf}) = 0.8823 \ (\pm 0.0378) \log k_w^{'}(\text{ff}) + 0.1474 \ (\pm 0.0654)$$
 (Eq. 3) ("sf" is standard flow at 1 ml/minute and "ff" is fast flow at 1.5 ml/minute)

N = 56, $R^2 = 0.910$, R = 0.954, s = 0.175, F = 544, $q^2 = 0.9$

Similarly, we have obtained good results by increasing the flow rate of the high lipophilicity range, from 2 to 3 mL/min (data not shown).

As a further caveat with the use of shake-methods we would like to report the widely different results we obtained with guanoxan (a guanidine derivative, CAS no. 5714-04-5) for which a value of -0.83 was reported as $logD_{oct}$. Using the shake-vial procedure B (see Experimental Section) values of -0.1 and -0.3 were obtained, in duplicate determinations. In a fairly extensive $logD_{oct}$ vs. concentration study, using the shake-vial procedure A, and we found that the values ranged from -1.6 to -1.0 upon decreasing the concentration, from 1.5 to 0.1 mg, in a 50:2 octanol:buffer system. Indeed Perlman²³ has reported a large variation in the $logD_{oct}$ values of diarylguanidines, up to $2 logD_{oct}$ units depending upon the counterion present, and that might be the case here. Under our conditions we found an $logD_{oct}$ value of -0.3, which is in very close agreement with the data from shake-flask procedure B but deviate significantly from the values from procedure A, even at the lowest concentration we have reached, and might be borderline acceptable for an estimation, against the literature data.

Most the solutes were purchased directly from commercial sources (Aldrich, Fluka, ICN, RBI, Sigma, Tocris) and used as received, in all cases. In several cases they were available as proprietary compounds or samples extracted from commercial formulations. Deionized water, HPLC grade methanol (J. P. Baker) and 1-octanol (Fisher Scientific) were used throughout.

The mobile phase consisted, in all cases, of 20 mM MOPS buffer at pH 7.4, with the addition of 0.15% of n-decylamine, ^{12,13} and methanol in varying proportions from 70 to 15% v/v. A 0.25% (v/v) amount of octanol was added to methanol, and octanol-saturated water was used to prepare the buffer. The mobile phase is now routinely prepared, by Brand-Nu Labs, Meriden, CT.

The capacity factors data ($k' = (t_R - t_0)/t_0$), obtained at various amounts of methanol, $\log k'$ values were then extrapolated to 0% methanol and reported as $\log k'_w$, using a linear procedure. In all cases, except for allopurinol ($R^2 = 0.96$), the square of the correlation coefficient was ≥ 0.99 . Injections of pure methanol were used to determine t_0 , i.e. the dead

time, while t_R has the usual meaning of the retention time for the analyte. For very low logD_{oct} compounds, atenolol (CAS no. 29122-68-7) was used to determine to, and is now used routinely for the low range.

All the chromatographic runs were performed on an Agilent 1100 HPLC ChemStation at the ambient temperature. The HPLC column used was Supelcosil LC-ABZ, 5 µm, 4.6 x 50 mm. A diode array detector was used to monitor signals at 235, 255, 265, 275 and 310 nm. We also tested columns manufactured from different silica bond lots, to ensure reproducibility. Samples were dissolved in 1:1 methanol/water in a concentration range of 50-100 µg/mL. The flow rate was 0.5, 1 or 2 mL/min, depending on the lipophilicity range. Three experimental lipophilicity ranges were established using, in all cases, three points for the extrapolation to

 $\log k_w$, as described as follows:

ElogD _{oct} Range	Flow rate (mL/min)	% MeOH	
< 1	0.5	15, 20, 25	
1-3	1	40, 45, 50	
> 3	2	60, 65, 70	

The samples are placed in the appropriate range by estimating their lipophilicity via computed values, or by prior experience with a given class. Experimental values obtained from runs outside the appropriate range, are typically run de novo, but a "screen" using a single injection at 75% methanol can be performed, to "weed out" high lipophilicity compounds. If the retention time is > 1.1 minutes, at a flow rate of 2 mL/min, the compound would likely yield an ElogDoct > 5. The user may then decide to adjust the conditions for that compound, such as the duration of each run, or to use such an estimated value, thus increasing the throughput and guarding against potential carry-overs.

In each case the entire group of samples is run before the column is equilibrated to the next condition, in an automated fashion. We typically start from the high range (highest methanol content) and run, sequentially, all the compounds. For the low range it was found that a period of equilibration between 1.5 to 2 hours is needed. At the end of a complete run the column is flushed with acetonitrile, at 2.0 mL/min, for 10-20 minutes.

The data analysis is then performed via an automated procedure relying on in-house software, which yields the ElogDoct values directly from the chromatographic data files.

The shake-flask logD_{oct} data, and in some cases data from countercurrent chromatography, were taken from the literature, after careful evaluation of the experimental method and temperature reported (generally between 20 and 25 °C) in the original references, or they were determined, except when data were not available or could not be determined experimentally due to the high lipophilicity of the compound. In such cases (clotrimazole and tolnaftate) a computed value was used. The shake-vial experimental

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measurements performed (Procedure A) were all conducted at least in duplicate, in amber glass vials and, in some cases, with varying ratios of octanol and MOPS buffer, always mutually pre-saturated prior to the experiment. The vials were shaken at least overnight. HPLC analysis at different wavelengths, after centrifugation and separation of the phases, was used for the quantitative analysis, and both phases were analyzed. In some cases compounds were used in a semi-automated logDoct shake-vial determination, using a phosphate buffer at pH 7.4 as the aqueous phase. In this case (Procedure B) a 1:1 ratio of noctanol and buffer (both phases were mutually pre-saturated) was used, with an agitation time ≥ 30 minutes, followed by centrifugation and analysis of both phases.

In several instances, as indicated in Table 1, the data were obtained from pH-metric determinations.

Statistical Analysis

All regression analyses were performed via the JMP statistical software package (v. 3.2.1, SAS Institute Inc.). Ten compounds were selected across the set of 90 compounds, covering the entire range of lipophilicity, to monitor the day-to-day performance of the method. Statistical calculations showed that the use of the 10 compounds would assure that the estimated slope, in the final regression equation, would be within \pm 0.09 of true one. The JMP software was also used for the Quality Monitoring. Data accumulated for the standard set of compounds, and regularly plotted on the control charts, constitute a powerful method for the detection of trends and variations in performance. Variations in $\log k_w$ values, for the selected compounds, should not exceed \pm 3 k_s, where k_s is the standard deviation estimate based on data collected under well controlled experiments. Triflupromazine $\log k_w$ are plotted for every run, to ensure that the column is performing suitably.

Table 1. Retention time and logD_{oct} data for the 90 solutes used.

Compound	CAS no.	$\log k_w^{'}$	s.d. ^c	ElogD _{oct}	logD _{oct} e	Res.	Refs.
Acebutolol	37517-30-9	-0.53	0.04	-0.39	-0.29	0.10	25,26
Acetominophen	103-90-2	0.15	0.01	0.38	0.51	0.13	27
Acetophenone	98-86-2	1.21	0.03	1.57	1.58	0.01	28
Allopurinol*	315-30-0	-0.27	0.01	-0.10	-0.44	-0.34	g, 29
Alprazolam	28981-97-7	1.73	0.04	2.16	2.12	-0.04	30
Alprenolol	13655-52-2	0.37	0.04	0.62	0.97	0.35	j,25
Amiodarone	1951-25-3	5.10	0.21	5.95	6.10	0.15	31
Amlodipine	88150-42-9	1.72	0.05	2.15	1.66	-0.49	g,h,2,32
Antipyrine	60-80-0	0.12	0.03	0.34	0.38	0.04	33
Atropine	51-55-8	-0.33	0.19	-0.16	-0.55	-0.39	j,34
Bifonazole	60628-96-8	4.33	0.07	5.09	4.77	-0.32	35
Bromazepam	1812-30-2	1.04	0.04	1.38	1.65	0.27	36,37
3-Bromoquinoline	5332-24-1	2.53	0.06	3.06	3.03	-0.03	38
Caffeine	58-08-2	-0.19	0.02	-0.01	-0.07	-0.06	35
Carbamazepine	298-46-4	1.40	0.03	1.79	2.19	0.41	39
Chloramphenicol*	56-75-7	1.19	0.07	1.55	1.14	-0.41	35
3-Chlorophenol	108-43-0	2.58	0.01	3.11	2.50	-0.61	28
Chlorpheniramine	132-22-9	1.20	0.16	1.56	1.41	-0.15	j,18,40
Chlorpromazine	50-53-3	2.66	0.08	3.20	3.38	0.18	18,26,4
Chlorthalidone	77-36-1	0.76	0.10	1.06	1.11	0.05	g,j,42
Cimetidine	51481-61-9	0.17	0.03	0.40	0.35	-0.05	j
Clonidine	4205-90-7	0.07	0.03	0.29	0.62	0.33	43
Clotrimazole	23593-75-1	4.03	0.06	4.75	5.20	0.45	16
Clozapine	5786-21-0	2.82	0.03	3.38	3.13	-0.25	j,18
Cocaine	50-36-2	0.24	0.05	0.48	1.05	0.57	44
Codeine	76-57-3	0.16	0.13	0.39	0.23	-0.16	41,45
Cyclothiazide	2259-96-3	2.15	0.09	2.63	2.09	-0.54	j
Deprenyl	2323-36-6	2.19	0.11	2.67	2.70	0.03	31
Desipramine	50-47-5	0.97	0.06	1.30	1.28	-0.02	g,j,46
Dexamethasone	50-02-2	1.62	0.04	2.03	1.83	-0.20	47
Diazepam	439-14-5	2.46	0.03	2.98	2.79	-0.19	48
3,5-	591-35-5	3.44	0.08	4.08	3.68	-0.40	38
Dichlorophenol							
Diethylstilbestrol	56-53-1	4.12	0.10	4.85	5.07	0.22	35

Compound	CAS no.	$\log k_w^{'b}$	s.d. ^c	ElogD _{oct}	logD _{oct} e	Res.	Refs.
Diltiazem	33286-22-5	1.59	0.22	2.00	2.06	0.06	
Diphenhydramine	58-73-1	1.04	0.06	1.38	1.29	-0.09	g,j,40
Disopyramide	3737-09-5	-1.21	0.10	-1.15	-0.66	0.49	j
Estradiol	50-28-2	3.28	0.08	3.90	4.01	0.11	35
Fentanyl citrate	990-73-8	1.94	0.09	2.39	2.91	0.52	49
Flecainide	54143-55-4	0.25	0.06	0.49	0.97	0.48	g,50
Fluconazole	86386-73-4	0.40	0.15	0.66	0.50	-0.16	51
Griseofulvin	126-07-8	1.73	0.06	2.16	2.18	0.02	35 '
Haloperidol	52-86-8	2.00	0.05	2.46	2.98	0.52	g,j
Hydrocortisone	50-23-7	1.14	0.06	1.49	1.55	0.06	49
Hydrocortisone-21	50-03-3	1.64	0.01	2.06	2.19	0.13	49
acetate							
Imipramine	50-49-7	1.56	0.19	1.97	2.40	0.43	g,j,18,40
Lidocaine	137-58-6	0.96	0.13	1.29	1.71	0.42	g,18,41
Loratadine	79794-75-5	4.02	0.07	4.73	4.44	-0.34	40
Lorazepam	846-49-1	2.30	0.03	2.80	2.51	-0.29	52
Lormetazepam	848-75-9	2.27	0.04	2.77	2.72	-0.05	53
Methotrimeprazine	60-99-1	2.10	0.27	2.57	2.77	0.20	j
Methylthioinosine	342-69-8	0.25	0.02	0.49	0.09	-0.40	16
Metoclopramide	364-62-5	0.46	0.16	0.73	0.64	-0.09	j
Metoprolol	56392-17-7	-0.73	0.08	-0.62	-0.16	0.46	25,31,34
							,46
Metronidazole	443-48-1	-0.08	0.02	0.12	-0.02	-0.14	54
Mexiletine .	31828-71-4	0.02	0.03	0.23	0.47	0.24	j
Morphine sulfate	64-31-3	0.10	0.10	0.32	0.03	-0.29	41,44,45
Naphthalene	91-20-3	3.03	0.06	3.62	3.37	-0.25	38
Nicotine	54-11-5	0.02	0.06	0.23	0.40	0.17	g,j
Nifedipine	21829-25-4	2.34	0.06	2.84	3.17	0.33	16
Nifuroxime	6236-05-1	1.14	0.07	1.49	1.28	-0.21	16
Nitrofurazone	59-87 - 0	0.29	0.01	0.53	0.23	-0.30	55
Nizatidine	76963-41-2	-0.13	0.04	0.06	-0.52	-0.58	56
Omeprazole	73590-58-6	1.59	0.04	2.00	2.30	0.30	g,57
Pentoxifylline	6493-05-6	0.03	0.02	0.24	0.29	0.05	58
Pirenzepine	28797-61-7	-0.15	0.06	0.04	-0.61	-0.65	g
Prednisolone	50-24-8	1.24	0.05	1.60	1.60	-0.00	16
Prednisone	53-03-2	0.90	0.06	1.22	1.46	0.24	35

Compound ^a	CAS no.	$\log k_w^{'}$	s.d. ^c	ElogD _{oct} ^d	logD _{oct} e	Res.	Refs.
Procainamide	51-06-9	-0.69	0.24	-0.57	-0.91	-0.34	59
Propafenone	54063-53-5	1.14	0.10	1.49	1.81	0.32	g,j
Propranolol	525-66-6	0.64	0.01	0.93	1.26	0.33	g,25,26,
							31,34,
							46,60
Quinidine	56-54-2	1.16	0.12	1.51	2.04	0.53	18,26,61
Quinoline	91-22-5	1.52	0.04	1.92	2.03	0.11	62
Ranitidine	66357-35-5	-0.63	0.01	-0.50	-0.29	0.21	j
Risperidone	106266-06-2	1.23	0.13	1.59	2.04	0.45	g
Sotalol	3930-20-9	-1.47	0.10	-1.45	-1.35	0.10	25,31
Sumatriptan	103628-46-2	-0.54	0.02	-0.40	- 1.00	-0.60	g,h,63
Terbutaline sulfate	23031-32-5	-1.51	0.06	-1.49	-1.35	0.14	31,46,64
Testosterone	58-22-0	2.63	0.04	3.17	3.29	0.12	47
Tetracaine	94-24-6	1.70	0.07	2.12	2.29	0.17	31
Thiamphenicol	15318-45-3	-0.05	0.01	0.15	-0.27	-0.42	65
Thioridazine	50-52-2	2.88	0.10	3.45	3.34	-0.11	j,66
Tiapride	51012-32-9	-0.58	0.05	-0.45	-0.90	-0.45	j
Tiotidine	69014-14-8	0.57	0.01	0.85	0.57	-0.28	g
Tolnaftate	2398-96-1	4.53	0.10	5.31	5.40	0.09	16
Trazodone	19794-93-5	2.45	0.06	2.97	2.54	-0.43	j
Triamterene	396-01-0	0.71	0.05	1.01	1.21	0.20	g,j,42,67
Trichlormethiazide	133-67-5	0.26	0.02	0.50	0.43	-0.07	g,j
Triflupromazine*	146-54-3	3.05	0.13	3.64	3.61	-0.03	j,66
Trimethoprim	738-70-5	0.36	0.02	0.61	0.63	0.02	j
Zaltidine	85604-00-8	0.53	0.02	0.80	0.74	-0.06	g

a. Asterisk denotes standard compound. b. Average of three to five determinations c.

Standard deviation of 3 to $5 \log k_w$ determinations. d. Equation 1. e. Average $\log D_{oct}$ from all the references or methods indicated. f. $\log D_{oct} - E \log D_{oct}$. g. Shake-vial Procedure A. h. Shake-vial Procedure B. j. Potentiometric determination.

5 Table 2. Retention time and logD_{oct} data for the 10 proprietary compounds in the test set.

Compound	$\log k_w^{'a}$	s.d. ^b	ElogD _{oct} c	logD _{oct} ^d	Residuals ^e
1	0.15	0.07	0.38	0.31	-0.07
2	0.79	0.03	1.10	1.16	0.06

Compound	$\log k_w^{'a}$	s.d. ^b	ElogD _{oct} ^c	logD _{oct} ^d	Residuals ^e
3	2.75	0.25	3.30	3.20	-0.10
4	0.30	0.06	0.55	0.68	0.13
5	1.14	0.12	1.49	1.66	0.17
6	2.17	0.10	2.65	2.19	-0.46
7	3.46	0.02	4.10	3.37	-0.73
8	2.70	0.01	3.25	2.58	-0.67
9	3.68	0.16	4.35	3.85	-0.50
10	2.14	0.06	2.62	2.10	-0.52
a.	Average of	three to five	determinations.	b. Standard	d deviation o

 $\log k_w^{'}$ determinations. c. Predicted ElogD_{oct} from Equation 1. d. Data from shake-vial and/or potentiometric determinations. The average was taken if more than one value was available. e. Average residual -0.27. ElogD_{oct} vs. logD_{oct}: $R^2 = 0.969$.

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